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Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
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Reviewer: Durreshwar Anjum

Timestamp: [year=2009; month=9; day=14; hr=15; min=28; sec=3; ms=286;]

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Application No: 10561867 Version No: 4.0

Input Set:

Output Set:

Started: 2009-09-10 16:40:27.577
Finished: 2009-09-10 16:40:27.938
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 361 ms
Total Warnings: 4
Total Errors: 0
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

| Error code | Error Description |
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| W 213 | Artificial or Unknown found in <213> in SEQ ID (6) |

SEQUENCE LISTING

<110> BURNOUF, Dominique, Yves, Joel
 WAGNER, Jerome, Edouard
 DUMAS, Philippe
 FUJII, Shingo
 FUCHS, Robert, Pierre, Paul
 OLIERIC, Vincent

<120> PROTEIN CRYSTAL COMPRISING THE PROCESSIVITY CLAMP FACTOR
 OF DNA POLYMERASE AND A LIGAND, AND ITS USES

<130> 0508-1147

<140> 10561867

<141> 2009-09-10

<150> PCT/EP2004/006942

<151> 2004-06-25

<150> EP 03291596.9

<151> 2003-06-27

<160> 6

<170> PatentIn version 3.5

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<211> 16

<212> PRT

<213> Escherichia coli

<400> 1

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| Val | Thr | Leu | Leu | Asp | Pro | Gln | Met | Glu | Arg | Gln | Leu | Val | Leu | Gly | Leu |
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<223> synthetic peptide

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| Arg | Pro | Val | Lys | Val | Thr | Pro | Asn | Gly | Ala | Glu | Asp | Glu | Ser | Ala | Glu |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |

| | | | | | |
|-----|-----|-----|-----|-----|-----|
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 <220>
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Val Ser Gly Pro Leu Gly Gly Arg Pro Thr Leu Pro Ile Leu Gly Asn
 20 25 30

Leu Leu Leu Gln Val Ala Asp Gly Thr Leu Ser Leu Thr Gly Thr Asp
 35 40 45

Leu Glu Met Glu Met Val Ala Arg Val Ala Leu Val Gln Pro His Glu
 50 55 60

Pro Gly Ala Thr Thr Val Pro Ala Arg Lys Phe Phe Asp Ile Cys Arg
 65 70 75 80

Gly Leu Pro Glu Gly Ala Glu Ile Ala Val Gln Leu Glu Gly Glu Arg
 85 90 95

Met Leu Val Arg Ser Gly Arg Ser Arg Phe Ser Leu Ser Thr Leu Pro
100 105 110

Ala Ala Asp Phe Pro Asn Leu Asp Asp Trp Gln Ser Glu Val Glu Phe
115 120 125

Thr Leu Pro Gln Ala Thr Met Lys Arg Leu Ile Glu Ala Thr Gln Phe
130 135 140

Ser Met Ala His Gln Asp Val Arg Tyr Tyr Leu Asn Gly Met Leu Phe
145 150 155 160

Glu Thr Glu Gly Glu Glu Leu Arg Thr Val Ala Thr Asp Gly His Arg
165 170 175

Leu Ala Val Cys Ser Met Pro Ile Gly Gln Ser Leu Pro Ser His Ser
180 185 190

Val Ile Val Pro Arg Lys Gly Val Ile Glu Leu Met Arg Met Leu Asp
195 200 205

Gly Gly Asp Asn Pro Leu Arg Val Gln Ile Gly Ser Asn Asn Ile Arg
210 215 220

Ala His Val Gly Asp Phe Ile Phe Thr Ser Lys Leu Val Asp Gly Arg
225 230 235 240

Phe Pro Asp Tyr Arg Arg Val Leu Pro Lys Asn Pro Asp Lys His Leu
245 250 255

Glu Ala Gly Cys Asp Leu Leu Lys Gln Ala Phe Ala Arg Ala Ala Ile
260 265 270

Leu Ser Asn Glu Lys Phe Arg Gly Val Arg Leu Tyr Val Ser Glu Asn
275 280 285

Gln Leu Lys Ile Thr Ala Asn Asn Pro Glu Gln Glu Glu Ala Glu Glu
290 295 300

Ile Leu Asp Val Thr Tyr Ser Gly Ala Glu Met Glu Ile Gly Phe Asn
305 310 315 320

Val Ser Tyr Val Leu Asp Val Leu Asn Ala Leu Lys Cys Glu Asn Val

325

330

335

Arg Met Met Leu Thr Asp Ser Val Ser Ser Val Gln Ile Glu Asp Ala
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Ala Ser Gln Ser Ala Ala Tyr Val Val Met Pro Met Arg Leu
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<213> Artificial Sequence

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<223> synthetic peptide - derived from SEQ ID NO 1

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